SEQUENCE LISTING

<110> Connex-Gesellschaft zur Optimerung von Forschung und Entwicklung mbH
INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE (I.N.S.E.R.M)

<120> Anti Hepatitis C virus antibody and uses therof

<130> b3030pct

<140> 09/744,176
<141> 2001-07-20

<150> ep 98 11 35 95.7
<151> 1998-07-21

<160> 22

<170> PatentIn version 3.2

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<220>
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acg gcc agg atc acc tgc tct gga gat gca ttg cca aag caa tat gct       96
Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala
            20                  25                  30

tac tgg tat cag cag aag cca ggc cag gcc cct gtg ttg gtg ata tat      144
Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35                  40                  45

aaa gat aat gag agg ccc tca ggg atc cct gag cga ttc tct ggc tcc      192
Lys Asp Asn Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50                  55                  60

agg tca ggg aca aca gtc acg ttg acc atc agt gga gtc cag gca gaa      240
Arg Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
65                  70                  75                  80

gac gag gct gac tat tac tgt caa tca gca gac agc agt ggt tct tcc      288
Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Ser Ser
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20 25 30

Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
35 40 45

Lys Asp Asn Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
50 55 60

Arg Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
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Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Ser Ser
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Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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1 5 10 15

acc ctg tcc ctc acc tgc gct gtc tat ggt ggg tcc tta agt ggt tac 96
Thr Leu Ser Leu Thr Cys Ala Val Tyr Gly Gly Ser Leu Ser Gly Tyr
20 25 30

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ttc tgg acc tgg atc cgc cag tcc ccc ggg aag ggg ctg gag tgg att      144
Phe Trp Thr Trp Ile Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Ile
        35                  40                  45

ggg gaa agc aat tat agt gga agt acc agg tac aac ccg tcc ctc aag      192
Gly Glu Ser Asn Tyr Ser Gly Ser Thr Arg Tyr Asn Pro Ser Leu Lys
    50                  55                  60

agt cga gtc acc ata tca gta gac acg tcc cag aac cag ttc tcc ctg      240
Ser Arg Val Thr Ile Ser Val Asp Thr Ser Gln Asn Gln Phe Ser Leu
65                  70                  75                  80

aag ctg agc tct gtg acc gcc gcg gac acg gct gta tat tac tgt gcg      288
Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala
                85                  90                  95

aga ggt tgg gcg gtg gac ggt atg gac gtc tgg ggc caa ggg acc acg      336
Arg Gly Trp Ala Val Asp Gly Met Asp Val Trp Gly Gln Gly Thr Thr
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Phe Trp Thr Trp Ile Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Ile
        35                  40                  45

Gly Glu Ser Asn Tyr Ser Gly Ser Thr Arg Tyr Asn Pro Ser Leu Lys
    50                  55                  60

Ser Arg Val Thr Ile Ser Val Asp Thr Ser Gln Asn Gln Phe Ser Leu
65                  70                  75                  80

Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala
                85                  90                  95
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Arg Gly Trp Ala Val Asp Gly Met Asp Val Trp Gly Gln Gly Thr Thr
            100                 105                 110

Val Thr Val Ser Ser
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tcc tct gag ctg aca cag cca ccc tcg gtg tca gtg tcc cca gga cag       48
Ser Ser Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1               5                   10                  15

acg gcc agg atc acc tgc tct gga gat gca ttg cca aag caa tat gct       96
Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala
            20                  25                  30

tac tgg tat cag cag aag cca ggc cag gcc cct gtg ttg gtg ata tat      144
Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
        35                  40                  45

aaa gat aat gag agg ccc tca ggg atc cct gag cga ttc tct ggc tcc      192
Lys Asp Asn Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
    50                  55                  60

agg tca ggg aca aca gtc acg ttg acc atc agt gga gtc cag gca gaa      240
Arg Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
65                  70                  75                  80

gac gag gct gac tat tac tgt caa tca gca gac agc agt ggt tct tcc      288
Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Ser Ser
                85                  90                  95

tgg gtg ttc ggc gga ggg acc aag ctg acc gtc cta                      324
Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105
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Ser Ser Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
1               5                   10                  15
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Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala
20 25 30

Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
35 40 45

Lys Asp Asn Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
50 55 60

Arg Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
65 70 75 80

Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Ser Ser
85 90 95

Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
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<210> 7
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<210> 9
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<210> 20
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<212> DNA
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<220>
<223> 3'-VL Primer: CL2

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